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1627

RAW SEQUENCE LISTING

DATE: 03/05/2002

PATENT APPLICATION: US/09/424,482C

TIME: 13:08:30

#20

Input Set : N:\Crf3\02212002\I424482A.raw

Output Set: N:\CRF3\03052002\I424482C.raw

1 <110> APPLICANT: CHOO, YEN
 2 KLUG, AARON
 3 ISALAN, MARK
 4 <120> TITLE OF INVENTION: NUCLEIC ACID BINDING POLYPEPTIDE LIBRARY
 5 <130> FILE REFERENCE: 71278/264974/BET
 C--> 6 <140> CURRENT APPLICATION NUMBER: US/09/424,482C
 7 <141> CURRENT FILING DATE: 2000-06-29
 8 <150> PRIOR APPLICATION NUMBER: 09/424,482
 9 <151> PRIOR FILING DATE: 1999-11-23
 10 <150> PRIOR APPLICATION NUMBER: PCT/GB98/01510
 11 <151> PRIOR FILING DATE: 1998-05-26
 12 <150> PRIOR APPLICATION NUMBER: GB 9710809.6
 13 <151> PRIOR FILING DATE: 1997-05-23
 14 <160> NUMBER OF SEQ ID NOS: 114
 15 <170> SOFTWARE: PatentIn version 2.1
 17 <210> SEQ ID NO: 1
 18 <211> LENGTH: 9
 19 <212> TYPE: DNA
 20 <213> ORGANISM: Artificial Sequence
 21 <220> FEATURE:
 22 <223> OTHER INFORMATION: Description of Artificial Sequence: LIB-A DNA sorting
 23 sequence
 24 <221> NAME/KEY: modified_base
 25 <222> LOCATION: (2)
 26 <223> OTHER INFORMATION: a, t, c or g
 27 <221> NAME/KEY: modified_base
 28 <222> LOCATION: (3)..(4)
 29 <223> OTHER INFORMATION: variable nucleotide
 30 <400> SEQUENCE: 1
 W--> 31 gnnncggcg 9
 33 <210> SEQ ID NO: 2
 34 <211> LENGTH: 9
 35 <212> TYPE: DNA
 36 <213> ORGANISM: Artificial Sequence
 37 <220> FEATURE:
 38 <223> OTHER INFORMATION: Description of Artificial Sequence: LIB-B DNA sorting
 39 sequence
 40 <221> NAME/KEY: modified_base
 41 <222> LOCATION: (3)..(4)
 42 <223> OTHER INFORMATION: variable nucleotide
 43 <400> SEQUENCE: 2
 W--> 44 gcnnccggcg 9
 46 <210> SEQ ID NO: 3

ENTERED

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47 <211> LENGTH: 9
48 <212> TYPE: DNA
49 <213> ORGANISM: Artificial Sequence
50 <220> FEATURE:
51 <223> OTHER INFORMATION: Description of Artificial DNA: LIB 1/2 sorting
52     sequence
53 <221> NAME/KEY: modified_base
54 <222> LOCATION: (5)..(9)
55 <223> OTHER INFORMATION: Any given nucleotide
56 <400> SEQUENCE: 3
W--> 57     gcggnnnnn
58 <210> SEQ ID NO: 4
59 <211> LENGTH: 31
60 <212> TYPE: PRT
61 <213> ORGANISM: Artificial Sequence
62 <220> FEATURE:
63 <223> OTHER INFORMATION: Description of Artificial Sequence: Zinc finger
64 <221> NAME/KEY: MOD_RES
65 <222> LOCATION: (1)..(2)
66 <223> OTHER INFORMATION: Any amino acid
67 <221> NAME/KEY: MOD_RES
68 <222> LOCATION: (4)..(8)
69 <223> OTHER INFORMATION: Any amino acid
70 <221> NAME/KEY: MOD_RES
71 <222> LOCATION: (10)..(23)
72 <223> OTHER INFORMATION: Any amino acid
73 <221> NAME/KEY: MOD_RES
74 <222> LOCATION: (25)..(30)
75 <223> OTHER INFORMATION: Any amino acid
76 <221> NAME/KEY: MOD_RES
77 <222> LOCATION: (31)
78 <223> OTHER INFORMATION: His or Cys
79 <223> OTHER INFORMATION: Positions 1-2 may vary in length from
80     0-2 residues
81 <223> OTHER INFORMATION: Positions 4-8 may vary in length from
82     1-5 residues
83 <223> OTHER INFORMATION: Positions 10-23 may vary in length from
84     9-14 residues
85 <223> OTHER INFORMATION: Positions 25-30 may vary in length from
86     3-6 residues
87 <400> SEQUENCE: 4
W--> 88     Xaa Xaa Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa
89         1             5             10             15
W--> 90     Xaa Xaa Xaa Xaa Xaa Xaa His Xaa Xaa Xaa Xaa Xaa Xaa Xaa
91         20             25             30
92 <210> SEQ ID NO: 5
93 <211> LENGTH: 24
94 <212> TYPE: PRT
95 <213> ORGANISM: Artificial Sequence

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98 <220> FEATURE:
99 <223> OTHER INFORMATION: Description of Artificial Sequence: Zinc finger
100 <221> NAME/KEY: MOD_RES
101 <222> LOCATION: (1)
102 <223> OTHER INFORMATION: Any amino acid
103 <221> NAME/KEY: MOD_RES
104 <222> LOCATION: (3)..(6)
105 <223> OTHER INFORMATION: Any amino acid
106 <221> NAME/KEY: MOD_RES
107 <222> LOCATION: (8)..(10)
108 <223> OTHER INFORMATION: Any amino acid
109 <221> NAME/KEY: MOD_RES
110 <222> LOCATION: (12)..(16)
111 <223> OTHER INFORMATION: Any amino acid
112 <221> NAME/KEY: MOD_RES
113 <222> LOCATION: (15)..(16)
114 <223> OTHER INFORMATION: Any amino acid
115 <221> NAME/KEY: MOD_RES
116 <222> LOCATION: (18)..(19)
117 <223> OTHER INFORMATION: Any amino acid
118 <221> NAME/KEY: MOD_RES
119 <222> LOCATION: (21)..(23)
120 <223> OTHER INFORMATION: Any amino acid
121 <223> OTHER INFORMATION: Positions 3-6 may vary in length from
122     2-4 residues
123 <223> OTHER INFORMATION: Positions 8-10 may vary in length from
124     2-3 residues
125 <400> SEQUENCE: 5
W--> 126     Xaa Cys Xaa Xaa Xaa Cys Xaa Xaa Xaa Phe Xaa Xaa Xaa Xaa Xaa
127         1             5             10             15
W--> 128     Leu Xaa Xaa His Xaa Xaa Xaa His
129         20
131 <210> SEQ ID NO: 6
132 <211> LENGTH: 4
133 <212> TYPE: PRT
134 <213> ORGANISM: Artificial Sequence
135 <220> FEATURE:
136 <223> OTHER INFORMATION: Description of Artificial Sequence: Linker
137 <400> SEQUENCE: 6
138     Thr Gly Glu Lys
139     1
141 <210> SEQ ID NO: 7
142 <211> LENGTH: 5
143 <212> TYPE: PRT
144 <213> ORGANISM: Artificial Sequence
145 <220> FEATURE:
146 <223> OTHER INFORMATION: Description of Artificial Sequence: Linker
147 <400> SEQUENCE: 7
148     Thr Gly Glu Lys Pro

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149      1      5
151 <210> SEQ ID NO: 8
152 <211> LENGTH: 26
153 <212> TYPE: PRT
154 <213> ORGANISM: Artificial Sequence
155 <220> FEATURE:
156 <223> OTHER INFORMATION: Description of Artificial Sequence: Consensus
157      structure sequence
158 <400> SEQUENCE: 8
159      Pro Tyr Lys Cys Pro Glu Cys Gly Lys Ser Phe Ser Gln Lys Ser Asp
160      1      5      10      15
161      Leu Val Lys His Gln Arg Thr His Thr Gly
162      20      25
164 <210> SEQ ID NO: 9
165 <211> LENGTH: 29
166 <212> TYPE: PRT
167 <213> ORGANISM: Artificial Sequence
168 <220> FEATURE:
169 <223> OTHER INFORMATION: Description of Artificial Sequence: Consensus
170      structure sequence
171 <400> SEQUENCE: 9
172      Pro Tyr Lys Cys Ser Glu Cys Gly Lys Ala Phe Ser Gln Lys Ser Asn
173      1      5      10      15
174      Leu Thr Arg His Gln Arg Ile His Thr Gly Glu Lys Pro
175      20      25
177 <210> SEQ ID NO: 10
178 <211> LENGTH: 6
179 <212> TYPE: PRT
180 <213> ORGANISM: Artificial Sequence
181 <220> FEATURE:
182 <223> OTHER INFORMATION: Description of Artificial Sequence: Leader peptide
183 <400> SEQUENCE: 10
184      Met Ala Glu Glu Lys Pro
185      1      5
187 <210> SEQ ID NO: 11
188 <211> LENGTH: 9
189 <212> TYPE: DNA
190 <213> ORGANISM: Artificial Sequence
191 <220> FEATURE:
192 <223> OTHER INFORMATION: Description of Artificial DNA: LIB 2/3 DNA
193      sorting sequence
194 <221> NAME/KEY: modified_base
195 <222> LOCATION: (1)..(5)
196 <223> OTHER INFORMATION: Any given nucleotide
197 <400> SEQUENCE: 11
W--> 198      nnnnnnggcg
200 <210> SEQ ID NO: 12
201 <211> LENGTH: 9
202 <212> TYPE: DNA

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Input Set : N:\Crf3\02212002\I424482A.raw

Output Set: N:\CRF3\03052002\I424482C.raw

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203 <213> ORGANISM: Artificial Sequence
204 <220> FEATURE:
205 <223> OTHER INFORMATION: Description of Artificial Sequence: Zinc finger -DNA
206     interaction sequence
207 <400> SEQUENCE: 12
208     cgcccacgc
209                                     9
210 <210> SEQ ID NO: 13
211 <211> LENGTH: 9
212 <212> TYPE: DNA
213 <213> ORGANISM: Artificial Sequence
214 <220> FEATURE:
215 <223> OTHER INFORMATION: Description of Artificial Sequence: Zinc finger-DNA
216     interaction sequence
217 <400> SEQUENCE: 13
218     acgcccacg
219                                     9
220 <210> SEQ ID NO: 14
221 <211> LENGTH: 9
222 <212> TYPE: DNA
223 <213> ORGANISM: Artificial Sequence
224 <220> FEATURE:
225 <223> OTHER INFORMATION: Description of Artificial Sequence: Zinc finger-DNA
226     interaction sequence
227 <400> SEQUENCE: 14
228     gcgtgggcg
229                                     9
230 <210> SEQ ID NO: 15
231 <211> LENGTH: 9
232 <212> TYPE: DNA
233 <213> ORGANISM: Artificial Sequence
234 <220> FEATURE:
235 <223> OTHER INFORMATION: Description of Artificial Sequence: Zinc finger-DNA
236     interaction library designed sequence
237 <221> NAME/KEY: modified_base
238 <222> LOCATION: (7)..(9)
239 <223> OTHER INFORMATION: Any given nucleotide
240 <400> SEQUENCE: 15
W--> 241     acgccgnnn
242                                     9
243 <210> SEQ ID NO: 16
244 <211> LENGTH: 36
245 <212> TYPE: PRT
246 <213> ORGANISM: Artificial Sequence
247 <220> FEATURE:
248 <223> OTHER INFORMATION: Description of Artificial Sequence: LIB-A and
249     LIB-B Zinc finger
250 <400> SEQUENCE: 16
251     Met Ala Glu Glu Arg Pro Tyr Ala Cys Pro Val Glu Ser Cys Asp Arg
252     1           5           10           15
253     Arg Phe Ser Arg Ser Asp Glu Leu Thr Arg His Ile Arg Ile His Thr
254     20           25           30
255     Gly Gln Lys Pro

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VERIFICATION SUMMARY

DATE: 03/05/2002

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Input Set : N:\Crf3\02212002\I424482A.raw

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L:6 M:270 C: Current Application Number differs, Wrong Format

L:31 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:44 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:57 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:89 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:91 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:126 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:128 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:241 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:271 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:287 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:303 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19